

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 09/870,884B  
Source: IFW/b  
Date Processed by STIC: 10/26/05

# ***ENTERED***



IFW16

## RAW SEQUENCE LISTING

DATE: 10/26/2005

PATENT APPLICATION: US/09/870,884B

TIME: 11:34:16

Input Set : A:\Sequence Listing new 10-20-05.txt

Output Set: N:\CRF4\10262005\I870884B.raw

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3 <110> APPLICANT: Hoeg-Jensen, Thomas
4   Havelund, Svend
5   Markussen, Jan
6   Ostergaard, Soren
7   Ridderberg, Signe
8   Balschmidt, Per
9   Schaffer, Lauge
10  Jonassen, Ib
12 <120> TITLE OF INVENTION: Glucose Dependent Release of Insulin from Glucose Sensing
Insulin
13   Derivatives
15 <130> FILE REFERENCE: 6213.200-US
17 <140> CURRENT APPLICATION NUMBER: 09/870,884B
18 <141> CURRENT FILING DATE: 2001-05-31
20 <150> PRIOR APPLICATION NUMBER: US 60/213,375
21 <151> PRIOR FILING DATE: 2000-06-23
23 <150> PRIOR APPLICATION NUMBER: Denmark PA 2000 00858
24 <151> PRIOR FILING DATE: 2000-06-02
26 <160> NUMBER OF SEQ ID NOS: 18
28 <170> SOFTWARE: PatentIn version 3.3
30 <210> SEQ ID NO: 1
31 <211> LENGTH: 21
32 <212> TYPE: PRT
33 <213> ORGANISM: Homo sapiens
35 <400> SEQUENCE: 1
37 Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
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41 Glu Asn Tyr Cys Asn
42           20
45 <210> SEQ ID NO: 2
46 <211> LENGTH: 30
47 <212> TYPE: PRT
48 <213> ORGANISM: Homo sapiens
50 <400> SEQUENCE: 2
52 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
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56 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr
57           20           25           30
60 <210> SEQ ID NO: 3
61 <211> LENGTH: 30
62 <212> TYPE: PRT
63 <213> ORGANISM: Homo sapiens
65 <400> SEQUENCE: 3
67 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr

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68 1           5           10           15
71 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Asp Lys Thr
72           20           25           30
75 <210> SEQ ID NO: 4
76 <211> LENGTH: 30
77 <212> TYPE: PRT
78 <213> ORGANISM: Homo sapiens
80 <400> SEQUENCE: 4
82 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
83 1           5           10           15
86 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Lys Pro Thr
87           20           25           30
90 <210> SEQ ID NO: 5
91 <211> LENGTH: 21
92 <212> TYPE: PRT
93 <213> ORGANISM: Homo sapiens
95 <400> SEQUENCE: 5
97 Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
98 1           5           10           15
101 Glu Asn Tyr Cys Gly
102           20
105 <210> SEQ ID NO: 6
106 <211> LENGTH: 30
107 <212> TYPE: PRT
108 <213> ORGANISM: Homo sapiens
110 <400> SEQUENCE: 6
112 Phe Val Lys Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
113 1           5           10           15
116 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Ile Lys Thr
117           20           25           30
120 <210> SEQ ID NO: 7
121 <211> LENGTH: 21
122 <212> TYPE: PRT
123 <213> ORGANISM: Homo sapiens
125 <400> SEQUENCE: 7
127 Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
128 1           5           10           15
131 Glu Asn Tyr Cys Asp
132           20
135 <210> SEQ ID NO: 8
136 <211> LENGTH: 29
137 <212> TYPE: PRT
138 <213> ORGANISM: Homo sapiens
140 <400> SEQUENCE: 8
142 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
143 1           5           10           15
146 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys
147           20           25
150 <210> SEQ ID NO: 9

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151 <211> LENGTH: 30
152 <212> TYPE: PRT
153 <213> ORGANISM: Homo sapiens
155 <400> SEQUENCE: 9
157 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
158 1 5 10 15
161 Leu Val Cys Gly Glu Arg Gly Phe Phe Phe Thr Pro Lys Thr
162 20 25 30
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166 <211> LENGTH: 30
167 <212> TYPE: PRT
168 <213> ORGANISM: Homo sapiens
171 <220> FEATURE:
172 <221> NAME/KEY: MISC_FEATURE
173 <222> LOCATION: (30)..(30)
174 <223> OTHER INFORMATION: x = ornithine
176 <400> SEQUENCE: 10
178 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
179 1 5 10 15
W--> 182 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Xaa
183 20 25 30
186 <210> SEQ ID NO: 11
187 <211> LENGTH: 30
188 <212> TYPE: PRT
189 <213> ORGANISM: Homo sapiens
192 <220> FEATURE:
193 <221> NAME/KEY: MISC_FEATURE
194 <222> LOCATION: (30)..(30)
195 <223> OTHER INFORMATION: x = diaminopropionic acid
197 <400> SEQUENCE: 11
199 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
200 1 5 10 15
W--> 203 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Xaa
204 20 25 30
207 <210> SEQ ID NO: 12
208 <211> LENGTH: 30
209 <212> TYPE: PRT
210 <213> ORGANISM: Homo sapiens
212 <400> SEQUENCE: 12
214 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
215 1 5 10 15
218 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Lys
219 20 25 30
222 <210> SEQ ID NO: 13
223 <211> LENGTH: 31
224 <212> TYPE: PRT
225 <213> ORGANISM: Homo sapiens
227 <400> SEQUENCE: 13
229 Pro Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu

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230 1           5           10           15
233 Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Thr
234           20           25           30
237 <210> SEQ ID NO: 14
238 <211> LENGTH: 30
239 <212> TYPE: PRT
240 <213> ORGANISM: Homo sapiens
242 <400> SEQUENCE: 14
244 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
245 1           5           10           15
248 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Asp
249           20           25           30
252 <210> SEQ ID NO: 15
253 <211> LENGTH: 30
254 <212> TYPE: PRT
255 <213> ORGANISM: Homo sapiens
257 <400> SEQUENCE: 15
259 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
260 1           5           10           15
263 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Glu
264           20           25           30
267 <210> SEQ ID NO: 16
268 <211> LENGTH: 30
269 <212> TYPE: PRT
270 <213> ORGANISM: Homo sapiens
273 <220> FEATURE:
274 <221> NAME/KEY: MISC_FEATURE
275 <222> LOCATION: (30)..(30)
276 <223> OTHER INFORMATION: x = O-aminoserine(BOC)
278 <400> SEQUENCE: 16
280 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
281 1           5           10           15
W--> 284 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Xaa
285           20           25           30
288 <210> SEQ ID NO: 17
289 <211> LENGTH: 30
290 <212> TYPE: PRT
291 <213> ORGANISM: Homo sapiens
294 <220> FEATURE:
295 <221> NAME/KEY: MISC_FEATURE
296 <222> LOCATION: (30)..(30)
297 <223> OTHER INFORMATION: x = diaminobutyric acid
299 <400> SEQUENCE: 17
301 Phe Val Asn Gln His Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr
302 1           5           10           15
W--> 305 Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr Pro Lys Xaa
306           20           25           30
309 <210> SEQ ID NO: 18
310 <211> LENGTH: 7

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Output Set: N:\CRF4\10262005\I870884B.raw

311 <212> TYPE: PRT  
312 <213> ORGANISM: homo sapien  
315 <220> FEATURE:  
316 <221> NAME/KEY: MISC\_FEATURE  
317 <222> LOCATION: (4)..(4)  
318 <223> OTHER INFORMATION: Xaa=NBPhe  
320 <220> FEATURE:  
321 <221> NAME/KEY: MISC\_FEATURE  
322 <222> LOCATION: (7)..(7)  
323 <223> OTHER INFORMATION: Xaa=Lys(lithocholoyl)  
325 <400> SEQUENCE: 18  
W--> 327 Gly Phe Phe Xaa Thr Pro Xaa  
328 1 5

RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 10/26/2005  
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Input Set : A:\Sequence Listing new 10-20-05.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:10; Xaa Pos. 30 //

Seq#:11; Xaa Pos. 30 //

Seq#:16; Xaa Pos. 30 //

Seq#:17; Xaa Pos. 30 //

Seq#:18; Xaa Pos. 4,7 //

## VERIFICATION SUMMARY

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Input Set : A:\Sequence Listing new 10-20-05.txt

Output Set: N:\CRF4\10262005\I870884B.raw

L:182 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:16  
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:16  
L:284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:16  
L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:16  
L:327 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0